

DATE: Tuesday, June 17, 2003

Set Name Query side by side			Set Name result set			
$DB=USPT,PGPB,JPAB,EPAB,DWPI;\ PLUR=YES;\ OP=OR$						
L13	L12 and (bladder or urinary).clm.	2	L13			
L12	L11 and (EGFR)	. 30	L12			
L11	(kDA or Dalton).clm.	5188	·L11			
L10	(urine and EGFR).clm.	5	L10			
L9	L7 and (EGF or EGFR or epidermal).ab.	. 4	L9			
L8	L7 and (EGF or EGFR or epidermal)	141	L8			
L7	13 and 16	298	L7			
L6	(37 or 37000) same (kda or dalton)	1925	L6			
L5	13 and (37 or 37000)	5715	L5			
L4	L3 and EGFR.ab.	. 9	L4			
L3	((424/9.1)!.CCLS. (435/7.1 435/7.2 435/7.21 435/7.22 435/7.23)!.CCLS. (436/64)!.CCLS.)	10953	L3			
DB=USPT; PLUR=YES; OP=OR						
L2	((435/7.1 435/7.2 435/7.21 435/7.22 435/7.23 435/436)!.CCLS.)	6374	L2			
$DB=USPT,PGPB,JPAB,EPAB,DWPI;\ PLUR=YES;\ OP=OR$						
L1	43	1506309	L1			

END OF SEARCH HISTORY

```
NEWS 30 Apr 11 Display formats in DGENE enhanced
                 MEDLINE Reload
 NEWS 31
         Apr 14
 NEWS 32
         Apr 17
                 Polymer searching in REGISTRY enhanced
 NEWS 33
         Jun 13
                 Indexing from 1947 to 1956 added to records in CA/CAPLUS
 NEWS 34 Apr 21
                 New current-awareness alert (SDI) frequency in
                  WPIDS/WPINDEX/WPIX
 NEWS 35
         Apr 28
                  RDISCLOSURE now available on STN
 NEWS 36
         May 05
                  Pharmacokinetic information and systematic chemical names
                  added to PHAR
 NEWS 37
         May 15
                 MEDLINE file segment of TOXCENTER reloaded
 NEWS 38
         May 15
                  Supporter information for ENCOMPPAT and ENCOMPLIT updated
 NEWS 39
         May 16
                 CHEMREACT will be removed from STN
         May 19
 NEWS 40
                 Simultaneous left and right truncation added to WSCA
 NEWS 41 May 19 RAPRA enhanced with new search field, simultaneous left and
                  right truncation
 NEWS 42
         Jun 06
                 Simultaneous left and right truncation added to CBNB
 NEWS 43
         Jun 06 PASCAL enhanced with additional data
 NEWS EXPRESS April 4 CURRENT WINDOWS VERSION IS V6.01a, CURRENT
               MACINTOSH VERSION IS V6.0b(ENG) AND V6.0Jb(JP),
               AND CURRENT DISCOVER FILE IS DATED 01 APRIL 2003
NEWS HOURS
              STN Operating Hours Plus Help Desk Availability
               General Internet Information
 NEWS INTER
NEWS LOGIN
              Welcome Banner and News Items
              Direct Dial and Telecommunication Network Access to STN
NEWS PHONE
NEWS WWW
              CAS World Wide Web Site (general information)
Enter NEWS followed by the item number or name to see news on that
```

specific topic.

All use of STN is subject to the provisions of the STN Customer agreement. Please note that this agreement limits use to scientific research. Use for software development or design or implementation of commercial gateways or other similar uses is prohibited and may result in loss of user privileges and other penalties.

* * * * * STN Columbus

FILE 'HOME' ENTERED AT 12:00:40 ON 17 JUN 2003

=> file .gary COST IN U.S. DOLLARS

SINCE FILE TOTAL ENTRY SESSION 0.21 0.21

FULL ESTIMATED COST

FILE 'MEDLINE' ENTERED AT 12:00:45 ON 17 JUN 2003

FILE 'CANCERLIT' ENTERED AT 12:00:45 ON 17 JUN 2003

FILE 'BIOSIS' ENTERED AT 12:00:45 ON 17 JUN 2003 COPYRIGHT (C) 2003 BIOLOGICAL ABSTRACTS INC. (R)

FILE 'EMBASE' ENTERED AT 12:00:45 ON 17 JUN 2003 COPYRIGHT (C) 2003 Elsevier Science B.V. All rights reserved.

FILE 'SCISEARCH' ENTERED AT 12:00:45 ON 17 JUN 2003 COPYRIGHT 2003 THOMSON ISI

=> s 37000

L1 320 37000

=> s l1 and antibod?

53 L1 AND ANTIBOD?

=> s 12 and (EGFR or epidermal)

=> s l1 and (Da or Daltons)

20 L1 AND (DA OR DALTONS)

=> dup rem 14

PROCESSING COMPLETED FOR L4

15 DUP REM L4 (5 DUPLICATES REMOVED)

=> d ibib abs 1-15

ANSWER 1 OF 15 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.

ACCESSION NUMBER: 1996:333791 BIOSIS

PREV199699056147 DOCUMENT NUMBER: TITLE:

Venom biochemical analysis as auxiliary tool to taxonomic distinction between Polybia (Myrapetra) paulista Ihering, 1896 and Polybia (Myrapetra) scutellaris (White, 1841),

(Hymenoptera, Vespidae.

Palma, Mario Sergio (1); Gobbi, Nivar; Manzoli-Palma, Maria AUTHOR (S):

De Fatima

CORPORATE SOURCE: (1) Dep. Biol., Centro Estudos Insetos Sociais, Inst.

Biociencias, UNESP - 13500 Rio Claro, SP Brazil

Revista Brasileira de Biologia, (1996) Vol. 56, No. 2, pp. SOURCE:

245-248.

ISSN: 0034-7108.

DOCUMENT TYPE: LANGUAGE:

Article

Portuguese

Portuguese; English SUMMARY LANGUAGE:

P. paulista and P. scutellaris presented respectively, 30 and 15 mu-g of proteins per sac of venom. The high performance gel permeation chromatography analysis of venoms revealed the presence of four peaks of proteins common to both species (MW 65000, 50000, 47500 and 27000 Daltons); four other peaks of MW 43000, 37000, 33000 and 31000 were in P. paulista venom. Peaks of MW 21500 and 1800 were present in the venom of P. scutellaris. The activity of some enzymes from crude venoms were assayed and revealed that P. paulista presented higher content of phospholipase A-2, hyaluronidase acid phosphatase and lipase activities than P. scutellaris. These results are sufficiently different to be used

as auxiliary tools for taxonomically separating P. paulista and P.

scutellaris.

ANSWER 2 OF 15 SCISEARCH COPYRIGHT 2003 THOMSON ISI L5

ACCESSION NUMBER: 95:152590 SCISEARCH

THE GENUINE ARTICLE: QH148

TITLE: ROSAT POINTED OBSERVATIONS OF 4 X-RAY BRIGHT DA

WHITE-DWARFS

AUTHOR: WOLFF B (Reprint); JORDAN S; BADE N; REIMERS D

CORPORATE SOURCE: CHRISTIAN ALBRECHTS UNIV KIEL, INST THEORET PHYS &

STENWARTE, D-24098 KIEL, GERMANY (Reprint); HAMBURGER STERNWARTE, D-21029 HAMBURG, GERMANY; CHRISTIAN ALBRECHTS

UNIV KIEL, STERNWARTE, D-24098 KIEL, GERMANY

COUNTRY OF AUTHOR: **GERMANY**

SOURCE: ASTRONOMY AND ASTROPHYSICS, (FEB 1995) Vol. 294, No. 1,

> pp. 183-189. ISSN: 0004-6361.

DOCUMENT TYPE:

Article; Journal PHYS

FILE SEGMENT:

ENGLISH

LANGUAGE: REFERENCE COUNT:

ABSTRACT IS AVAILABLE IN THE ALL AND IALL FORMATS

We present pointed observations of four hot DA white dwarfs with the ROSAT satellite. Measurements with the Position Sensitive Proportional Counter (PSPC) and the Wide Field Camera (WFC) were analyzed in the framework of pure hydrogen model atmospheres. The results fit well into the picture of previous studies that objects between about 20000 and 40000 K have almost pure hydrogen atmospheres, while at higher temperatures additional absorbers are present in general. Of particular interest are PG 0824+289, PG 1657+344, and PG 1658+441. The former is a close binary consisting of a DA white dwarf and a dwarf carbon star (dC). The effective temperature of 37000 K, derived from the ROSAT observations, is very close to the value determined from IUE spectra, which - unlike optical spectra - are not contaminated by the ''cool'' dC. Therefore we conclude that the DA component of the system has a pure hydrogen atmosphere. The magnetic DA PG 1658+441 has the highest mass (1.31 M(circle dot)) of all known single white dwarfs. Its X-ray fur can be explained by a pure hydrogen atmosphere. An optical spectrum of PG 1657+344 reveals that this object is one of the few DA white dwarfs with T-eff greater than or similar to 50000 K. In this case the analysis of the ROSAT data shows clear evidence for absorbing material besides hydrogen.

L5 ANSWER 3 OF 15 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.DUPLICATE

ACCESSION NUMBER: 1994:457489 BIOSIS DOCUMENT NUMBER: PREV199497470489

TITLE: Monoclonal antibodies for Streptomyces lividans and their

use for immunomagnetic capture of spores from soil.

AUTHOR(S): Wipat, Anil; Wellington, Elizabeth M. H.; Saunders, Venetia

A. (1)

CORPORATE SOURCE: (1) Sch. Biomolecular Sci., Liverpool John Moores

University, Liverpool L3 3AF UK

SOURCE: Microbiology (Reading), (1994) Vol. 140, No. 8, pp.

2067-2076.

DOCUMENT TYPE: Article LANGUAGE: English

Monoclonal antibodies were produced to Streptomyces lividans spore surface antigens. One particular hybridoma cell line, 43H6, produced a monoclonal antibody that reacted exclusively with Streptomyces cluster group 21 in an enzyme-linked immunosorbent assay (ELISA). Antibody 43H6 was found to be of subclass IgG1, kappa light chain. Western blot (immunoblot) analysis revealed that 43H6 recognized a major outer spore polypeptide of about 37000 Da. The epitope was stably maintained in S. lividans spores over at least seven sporulation cycles on laboratory medium and for at least 14 weeks in sterile soil systems. The species group specificity of antibody 43H6 was exploited in the development of an immunocapture technique for the isolation of streptomycetes from soil. Magnetic beads coated with antibody 43H6 were mixed with soil samples seeded with S. lividans spores. Spore-bead complexes were recovered using magnets. Treatment of beads with blocking agents and the inclusion of detergents in the recovery system lessened non-specific binding of spores to beads and improved recovery. In buffer solutions decreasing the spore concentration increased the recovery values for a fixed bead concentration. At a spore concentration of 5 times 10-7 ml-1 the recovery was 4.3% whilst at 5 times 10-2 ml-1 it was 76% for a fixed bead concentration of 0.6 mg ml-1. Using a bead concentration of 2 mg per 10 g soil, approximately 30% of the target spore population of 10-6 c.f.u. was recovered from sterile soil and 4% from non-sterile soil. This method offers a rapid means of selectively recovering and concentrating Streptomyces spores from soil samples.

5 ANSWER 4 OF 15 SCISEARCH COPYRIGHT 2003 THOMSON ISI

ACCESSION NUMBER: 94:531691 SCISEARCH

THE GENUINE ARTICLE: PC693

TITLE: MONOC

MONOCLONAL-ANTIBODIES FOR STREPTOMYCES-LIVIDANS AND THEIR

USE FOR IMMUNOMAGNETIC CAPTURE OF SPORES FROM SOIL

AUTHOR: WIPAT A; WELLINGTON E M H; SAUNDERS V A (Reprint)

CORPORATE SOURCE: LIVERPOOL JOHN MOORES UNIV, SCH BIOMOLEC SCI, LIVERPOOL L3

3AF, MERSEYSIDE, ENGLAND (Reprint); LIVERPOOL JOHN MOORES

UNIV, SCH BIOMOLEC SCI, LIVERPOOL L3

3AF, MERSEYSIDE,
ENGLAND; UNIV WARWICK, DEPT BIOL SCI, COVENTRY CV4 7AL, W

MIDLANDS, ENGLAND

COUNTRY OF AUTHOR:

ENGLAND

SOURCE:

MICROBIOLOGY-UK, (AUG 1994) Vol. 140, Part 8, pp.

2067-2076.

ISSN: 1350-0872.

DOCUMENT TYPE: FILE SEGMENT:

Article; Journal

LIFE ENGLISH

LANGUAGE:

33

REFERENCE COUNT:

ABSTRACT IS AVAILABLE IN THE ALL AND IALL FORMATS

Monoclonal antibodies were produced to Streptomyces lividans spore surface antigens. One particular hybridoma cell line, 43H6, produced a monoclonal antibody that reacted exclusively with Streptomyces cluster group 21 in an enzyme-linked immunosorbent assay (ELISA). Antibody 43H6 was found to be of subclass IgG1, kappa light chain. Western blot (immunoblot) analysis revealed that 43H6 recognized a major outer spore polypeptide of about 37000 Da. The epitope was stably maintained in S, lividans spores over at least seven sporulation cycles on laboratory medium and for at least 14 weeks in sterile soil systems. The species group specificity of antibody 43H6 was exploited in the development of an immunocapture technique for the isolation of streptomycetes from soil. Magnetic beads coated with antibody 43H6 were mixed with soil samples seeded with 5, lividans spores. Spore-bead complexes were recovered using magnets. Treatment of beads with blocking agents and the inclusion of detergents in the recovery system lessened non-specific binding of spores to beads and improved recovery. In buffer solutions decreasing the spore concentration increased the recovery values for a fixed bead concentration. At a spore concentration of $5 \times 10(7)$ ml(-1) the recovery was 4.3% whilst at 5 x 10(2) ml(-1) it was 76% for a fixed bead concentration of 0.6 mg ml(-1). Using a bead concentration of 2 mg per 10 g soil, approximately 30% of the target spore population of 10(6) c.f.u. was recovered from sterile soil and 4% from non-sterile soil. This method offers a rapid means of selectively recovering and concentrating Streptomyces spares from soil samples.

ANSWER 5 OF 15 MEDLINE DUPLICATE 2

ACCESSION NUMBER:

94264673 MEDLINE

DOCUMENT NUMBER:

94264673 PubMed ID: 8205112

TITLE: An improved method for the purification of human

erythropoietin with high in vivo activity from the urine of

anemic patients.

AUTHOR:

Inoue N; Wada M; Takeuchi M

CORPORATE SOURCE:

Pharmaceutical Laboratory, Kirin Brewery Co., Ltd., Gumma,

SOURCE:

BIOLOGICAL AND PHARMACEUTICAL BULLETIN, (1994 Feb) 17 (2)

180-4.

Journal code: 9311984. ISSN: 0918-6158.

PUB. COUNTRY:

Japan

DOCUMENT TYPE:

Journal; Article; (JOURNAL ARTICLE)

LANGUAGE:

English

FILE SEGMENT:

Priority Journals

ENTRY MONTH:

199407

ENTRY DATE:

Entered STN: 19940721

Last Updated on STN: 19980206 Entered Medline: 19940708

AB An improved method for the purification of human erythropoietin with high in vivo activity from urine was developed. This method involved ion-exchange, gel permeation, affinity chromatography, and reverse-phase chromatography but did not involve any stabilizing procedures. The purified human urinary erythropoietin showed a single broad band with a molecular weight between 37000 and 39000 Da on sodium dodecyl sulfate polyacrylamide gel electrophoresis, and had an in vivo specific activity of 160000 IU/mg comparable to that of human erythropoietin produced in recombinant Chinese hamster ovary cells. We found that omission of the phenol treatment and ethanol precipitation

which are usually used in the purification of human urinary erythropoietin greatly improved the biological activity of the final product. Phenol treatment followed by ethanol precipitation did not affect the amino acid composition but decreased the apparent molecular weight and N-acetylglucosamine content of human urinary erythropoietin. These findings suggest that phenol treatment followed by ethanol precipitation does not restore erythropoietin with high branched sugar chains which would have high in vivo specific activity as reported previously (M. Takeuchi, et al. (1989) Proc. Natl. Acad. Sci. U.S.A., 86, 7819-7822).

L5 ANSWER 6 OF 15 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.

ACCESSION NUMBER: 1992:12243 BIOSIS

DOCUMENT NUMBER: BA93:12243

TITLE: EXTRACELLULAR VACCINIA VIRUS FORMATION AND CELL-TO-CELL

VIRUS TRANSMISSION ARE PREVENTED BY DELETION OF THE GENE

ENCODING THE 37000-DALTON OUTER ENVELOPE PROTEIN.

AUTHOR(S): BLASCO R; MOSS B

CORPORATE SOURCE: LAB. VIRAL DIS., NATL. INST. ALLERGY INFECTIOUS DIS.,

BETHESDA, MD. 20892.

SOURCE: J VIROL, (1991) 65 (11), 5910-5920.

CODEN: JOVIAM. ISSN: 0022-538X.

FILE SEGMENT: BA; OLD LANGUAGE: English

There are two types of infectious vaccinia virus particles: intracellular naked virions and extracellular enveloped virions (EEV). To determine the biological role of the enveloped form of vaccinia virus, we produced and characterized a mutant that is defective in EEV formation. The strategy involved replacement by homologous recombination of the gene F13L, encoding a 37,000-Da protein (VP37) that is specific for the outer envelope of EEV, with a selectable antibiotic resistance marker, the Escherichia coli gpt gene. Initial experiments, however, suggested that such a mutation was lethal or prevented plaque formation. By employing a protocol consisting of high-multiplicity passages of intracellular virus from the transfected cells and then limiting dilution cloning, we succeeded in isolating the desired mutant, which was defective in production of plaques and extracellular virus but made normal amounts of intracellular naked virions. Electron microscopic examination indicated that the mutant virus particles, unlike wild type, were neither wrapped with Golgi-derived membranes nor associated with the cell surface. The absence of VP37 did not prevent the transport of the viral hemagglutinin to the plasma membrane but nevertheless abrogated both low-pH- and antibody-mediated cell fusion. These results indicate that VP37 is required for EEV formation and also plays a critical role in the local cell-to-cell transmission of vaccinia virus, perhaps via enveloped virions attached to or released from the cell membrane. By contrast, a mutated virus with a deletion of the K4L open reading frame, which is a homolog of the VP37 gene, was not defective in formation of plaques or EEV.

L5 ANSWER 7 OF 15 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.

ACCESSION NUMBER: 1991:55987 BIOSIS

DOCUMENT NUMBER: BA91:34268

TITLE: PROCESSING OF PSEUDOMONAS EXOTOXIN BY A CELLULAR PROTEASE

RESULTS IN THE GENERATION OF A 37000-DA

TOXIN FRAGMENT THAT IS TRANSLOCATED TO THE CYTOSOL.

AUTHOR(S): OGATA M; CHAUDHARY V K; PASTAN I; FITZGERALD D J

CORPORATE SOURCE: LAB. MOL. BIOL., NATIONAL CANCER INST., NATIONAL INST.

HEALTH, BETHESDA, MD. 20892.

SOURCE: J BIOL CHEM, (1990) 265 (33), 20678-20685.

CODEN: JBCHA3. ISSN: 0021-9258.

FILE SEGMENT: BA; OLD LANGUAGE: English

AB Pseudomonas exotoxin (PE) was incubated with cells and extracts analyzed for processed fragments. PE was proteolytically cleaved to produce a N-terminal 28-kDa and a C-terminal 37-kDa fragment, the latter being composed of a portion of domain II and all of domain III (the

ADP-ribosylating domain). Cleavage was evident at 10 min after toxin addition and endosome preparations contained the processed fragments. Initially, the two fragments were linked by a disulfide bond. Subsequently, the 37-kDa fragment was reduced and translocated to the cytosol where it inactivated protein synthesis. Cytosol from toxin-treated cells was greatly enriched in the 37-kDa fragment. The 37-kDa fragment appears to be essential for toxicity since mutant PE molecules that do not produce this fragment, or cannot deliver it to the cytosol, fail to kill cells.

L5 ANSWER 8 OF 15 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.

ACCESSION NUMBER: 1990:10308 BIOSIS

DOCUMENT NUMBER: BA89:10308

TITLE: ISOLATION AND CHARACTERIZATION OF THE NERVE GROWTH FACTOR

FROM THE VENOM OF THE MIDDLE ASIAN SNAKE

ECHIS-MULTISOUAMATUS.

AUTHOR(S): KHAMIDOV D KH; YUKEL'SON L YA; SALIKHOV R S; KHAFIZOVA M G

CORPORATE SOURCE: INST. BIOCHEM., ACAD. SCI. UZB. SSR, TASHKENT, USSR.

SOURCE: BIOKHIMIYA, (1989) 54 (6), 987-991.

CODEN: BIOHAO. ISSN: 0320-9725.

FILE SEGMENT: BA; OLD LANGUAGE: Russian

AB The nerve growth factor (NGF) was isolated from the Echis multisquamatus venom by ultrafiltration on PM-10 filter, chromatography on TSK-55 gel, ion-exchange chromatography on CM-cellulose and gel filtration on Sephadex G-75. The protein exhibited a marked nerve growth activity within the concentration range of 10-15 ng/ml in cultures of chicken embryo spinal ganglia. The molecular mass of NGF is equal to 33000-37000

Da according to Sephadex G-75 gel filtration data; however, according to SDS electrophoresis data its Mr is 13000 Da.

Isoelectrofocusing data suggest that the pI of the isolated factor lies in the region of 7.0-7.2; sugar content is 1-2%.

L5 ANSWER 9 OF 15 EMBASE COPYRIGHT 2003 ELSEVIER SCI. B.V.

ACCESSION NUMBER: 88265691 EMBASE

DOCUMENT NUMBER: 1988265691

TITLE: Heparin modulates the secretion of a major excreted

protein-like molecule by vascular smooth muscle cells.

AUTHOR: Cochran D.L.; Castellot Jr. J.J.; Robinson J.M.; Karnovsky

M.J.

CORPORATE SOURCE: Department of Periodontology, School of Dentistry, Virginia

Commonwealth University, Richmond, VA, United States

SOURCE: Biochimica et Biophysica Acta - General Subjects, (1988)

967/2 (289-295).

ISSN: 0304-4165 CODEN: BBGSB3

COUNTRY: Netherlands

DOCUMENT TYPE: Journal

FILE SEGMENT: 029 Clinical Biochemistry

030 Pharmacology

037 Drug Literature Index

LANGUAGE: English SUMMARY LANGUAGE: English

AB Previous work from our laboratory has shown that heparin specifically induces the release of a pair of proteins of approximately 35000 and 37000 Da into the culture medium of vascular smooth muscle cells (SMC). In this report, we demonstrate that the previously identified 37000-Da smooth muscle protein is composed of two protein species with very similar molecular weights based on migration patterns in SDS-polyacrylamide gels. The larger molecular weight species in this doublet has a similar molecular weight and shares antigenic determinants with major excreted protein (MEP), a lysosomal proteinase previously shown to be secreted by normal and transformed fibroblasts and epidermal cells. Antisera to MEP precipitated the higher molecular weight band from the doublet; preimmune serum was not reactive with the smooth muscle protein. Exposure of smooth muscle cells to heparin

resulted in decreased amounts of immunoprecipitable protein released into the medium. Thus, it now appears that three proteins in the 35000-38000 molecular weight range are modulated by heparin, and that the largest of the heparin-modulated vascular SMC) proteins has a similar molecular weight and is immunologically related to MEP. The release of MEP-like protein from SMC is decreased by heparin, while the remaining two heparin-modulated proteins are increased in the presence of heparin.

L5 ANSWER 10 OF 15 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.

ACCESSION NUMBER: 1987:167166 BIOSIS

DOCUMENT NUMBER: BA83:85607

TITLE: GENE ENCODING THE 37000-DALTON MINOR SIGMA FACTOR

OF BACILLUS-SUBTILIS RNA POLYMERASE ISOLATION NUCLEOTIDE

SEQUENCES CHROMOSOMAL LOCUS AND CRYPTIC FUNCTION.

AUTHOR(S): DUNCAN M L; KALMAN S S; THOMAS S M; PRICE C W

CORPORATE SOURCE: DEP. FOOD SCI. AND TECHNOL., UNIV. CALIFORNIA, DAVIS,

CALLE OF CLC

CALIF. 95616.

SOURCE: J BACTERIOL, (1987) 169 (2), 771-778.

CODEN: JOBAAY. ISSN: 0021-9193.

FILE SEGMENT: BA; OLD LANGUAGE: English

We began an analysis of rpoF, the gene encoding the cryptic, 37,000-dalton minor sigma factor (sigma-37) of Bacillus subtilis RNA polymerase. Using antibody raised against sigma-37 holoenzyme to probe a λ gt11 expression vector library, we isolated a 901-base-pair EcoRI fragment that expressed the COOH-terminal half of sigma-37 fused to lacZ. We used this fragment as a hybridization probe to isolate the entire rpoF gene and additional flanking sequences. Identity of the cloned gene was confirmed by the size and immunological reaction of its product expressed in Escherichia coli and, after DNA sequencing, by the homology of its predicted product (264 residues; 30,143 daltons) with other sigma factors. The DNA sequence also suggested that rpoF may lie in a gene cluster. Upstream of rpoF was an open reading frame that would encode a protein of 17,992 daltons; this frame overlapped the rpoF-coding sequence by 41 base pairs. Immediately following rpoF was a reading frame that would encode a protein of at least 20,000 daltons; expression of this region may be translationally coupled to that of rpoF. By plasmid integration and PBS1 transduction, we found the chromosomal locus of rpoF linked to ddl and dal at 40° on the B. subtilis map and near no known lesions affecting growth regulation or development. Further, an rpoF null mutation resulting from gene disruption had no effect on cell growth or sporulation in rich medium, suggesting that sigma-37 may partly control a regulon not directly involved in the sporulation process.

L5 ANSWER 11 OF 15 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.

ACCESSION NUMBER: 1986:379431 BIOSIS

DOCUMENT NUMBER: BA82:74407

TITLE: SIMILARITY BETWEEN GLYCERALDEHYDE-3-PHOSPHATE DEHYDROGENASE

AND A 37000-DALTON PROTEIN WHICH IS ABUNDANTLY

EXPRESSED IN HUMAN LUNG CANCERS.

AUTHOR(S): OHKUBO M; NAKAMURA Y; TOKUNAGA K; SAKIYAMA S

CORPORATE SOURCE: DIV. OF BIOCHEMISTRY, CHIBA CANCER CENT. RES. INST., 666-2,

NITONA-CHO, CHIBA 280.

SOURCE: JPN J CANCER RES (GANN), (1986) 77 (6), 554-559.

CODEN: JJCREP. ISSN: 0910-5050.

FILE SEGMENT: BA; OLD LANGUAGE: English

AB Human lung cancers of all histological types contain a protein of 37,000 daltons (37K) as an abundant component. Partial sequence analysis of purified 37K revealed a strong homology with glyceraldehyde-3-phosphate dehydrogenase (GAPDH, EC 1.2.1.12). Tryptic peptide mapping analysis showed that the pattern of 37K was very similar to those of GAPDHs both purified from lung tumor and obtained commercially. An antibody raised against 37K in a rabbit also reacted with authentic GAPDH. These results

suggest a possible involvement of GAPDH itself or a GAPDH-related protein in lung tumorigenesis.

L5 ANSWER 12 OF 15 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.

ACCESSION NUMBER: 1986:130479 BIOSIS

DOCUMENT NUMBER: BA81:40895

TITLE: EFFECTS OF ACTINOMYCIN D ANALOGS ON NUCLEOLAR

PHOSPHOPROTEIN B-23 37000 DALTONS

ISOELECTRIC POINT 5.1.

AUTHOR(S): YUNG B Y-M; BUSCH R K; BUSCH H; MAUGER A B; CHAN P-K CORPORATE SOURCE: BAYLOR COLL. MED., DEP. PHARMACOL., HOUSTON, TEX. 77030,

USA.

SOURCE: BIOCHEM PHARMACOL, (1985) 34 (22), 4059-4064.

CODEN: BCPCA6. ISSN: 0006-2952.

FILE SEGMENT: BA; OLD LANGUAGE: English

Localization of protein B23 in HeLa cells after treatment with actinomycin D and its analogs was studied using indirect immunofluorescence. Bright nucleolar fluorescence was observed in control HeLa cells. After treatment with actinomycin D (250 ng/ml) for 2 hr, a uniform nucleoplasmic fluorescence was observed. Similar results were obtained with the actinomyin analogs, actinomycin Z5 and actinomycin K2T. Only after a much longer incubation (24 hr) with actinomycin 4.4'-gly was nucleoplasmic fluorescence observed. Actinomycin D, actinomycin Z5, and actinomycin K2T inhibited [3H]uridine incorporation into the trichloroacetic acid insoluble fraction of HeLa cells with IC50 values of 9.5 ± 3.2, 59.1 \pm 19.6 and 1423.3 \pm 212.2 ng/ml respectively. No inhibition of [3H]uridine incorporation was observed using actinomycin 4.4'-gly (2000 ng/ml, 2-hr incubation). The order of potency for the loss of nucleolar fluorescence and the concurrent increase in nucleoplasmic fluorescence was actinomycin D > actinomycin Z5 > actinomycin K2T > actinomycin 4.4'-gly, which correlated with the order of their IC50 values for inhibition of [3H]uridine incorporation. Studies of the effects of actinomycin D and its analogs on RNA synthesis and localization of protein B23 indicated that there is a direct relationship between the B23 "translocation" from nucleolus to nucleoplasm and the inhibition of RNA synthesis. At 45-55% inhibition of RNA synthesis, both nuclear and nucleolar B23 immunofluorescence were observed. At 75-85% inhibition, only a uniform nucleoplasmic fluorescence was observed.

L5 ANSWER 13 OF 15 EMBASE COPYRIGHT 2003 ELSEVIER SCI. B.V.

ACCESSION NUMBER: 85178700 EMBASE

DOCUMENT NUMBER: 1985178700

TITLE: Purification and preliminary characterization of

2-monoacylglycerol acyltransferase from rat intestinal

villus cells.

AUTHOR: Manganaro F.; Kuksis A.

CORPORATE SOURCE: Banting and Best Department of Medical Research, University

of Toronto, Toronto, Ont. M5G 1L6, Canada

SOURCE: Canadian Journal of Biochemistry and Cell Biology, (1985)

63/5 (341-347). CODEN: CJBBDU

COUNTRY: Canada DOCUMENT TYPE: Journal

FILE SEGMENT: 029 Clinical Biochemistry

048 Gastroenterology

LANGUAGE: English SUMMARY LANGUAGE: French

AB We have purified the monoacylglycerol acyltransferase from rat small intestinal mucosa to homogeneity by a combination of hydrophobic absorption, guanidine dissociation, and gel filtration. The purified enzyme gives a single band of 37000 daltons on sodium dodecyl sulphate - polyacrylamide gel electrophoresis. The enzyme has a specific activity of about 5900 nmol/mg per hour and represents 0.12% of total cell protein, corresponding to about a 600-fold purification. The

enzyme does not acylate diacylglycerols to triacylglycerols, which is consistent with the separate physical existence of the mono- and di-acylglycerol acyltransferases. The enzyme acylates the 2-monoacylglycerols to yield an essentially racemic mixture of diacylglycerols. It does not acylate glycerol 3-phosphate.

L5 ANSWER 14 OF 15 EMBASE COPYRIGHT 2003 ELSEVIER SCI. B.V.

ACCESSION NUMBER: 82054122 EMBASE

DOCUMENT NUMBER: 1982054122

TITLE: Mode of degradation of myofibrillar proteins by an

endogenous protease, cathepsin L.

AUTHOR: Matsukura U.; Okitani A.; Nishimuro T.; Kato H.

CORPORATE SOURCE: Dept. Agric. Chem., Fac. Agric., Univ. Tokyo, Bunkyo-ku,

Tokyo, Japan

SOURCE: Biochimica et Biophysica Acta, (1981) 662/1 (41-47).

CODEN: BBACAQ

COUNTRY: Netherlands

DOCUMENT TYPE: Journal

FILE SEGMENT: 029 Clinical Biochemistry

LANGUAGE: English

The mode of degradation of myofibrils and their constituent proteins by cathepsin L (EC 3.4.22.15) of rabbit skeletal muscle was studied. Sodium dodecyl sulfate (SDS)-polyacrylamide gel electrophoresis showed that cathepsin L degraded myosin heavy chain, α-actinin, actin, troponin T and troponin I assembled in myofibrils and produced mainly fragments of 160000 and 30000 daltons in the acidic pH region. This degradation was most intense around pH 4. Degradation of myosin in the isolated state by cathepsin L resulted in the disappearance of the heavy chain and the decrease of light chains 1, 2 and 3, producing fragments of 160000, 92000, 83000 and 60000 daltons. The degradation of the heavy chain was most severe at pH 4.2. Cathepsin L degraded actin into fragments of 40000, 37000 and 30000 daltons. This action was most intense at pH 4.7. Tropomyosin was not degraded. Troponin T and troponin I were degraded into fragments of 30000 and 13000 daltons at pH 3.7-6.7, which were degraded further into smaller fragments. Troponin C was not degraded. α-Actinin was degraded into several fragments, the major one of which showed an M(r) of 80000. This degradation was most intense at pH 3.0-3.5.

L5 ANSWER 15 OF 15 EMBASE COPYRIGHT 2003 ELSEVIER SCI. B.V.

ACCESSION NUMBER: 74129174 EMBASE

DOCUMENT NUMBER: 1974129174

TITLE: The action of Factor Xa, thrombin and trypsin on human

Factor II.

AUTHOR: Kisiel W.; Hanahan D.J.

CORPORATE SOURCE: Dept. Biochem., Coll. Med., Univ. Arizona, Tucson, Ariz.

85724, United States

SOURCE: Biochimica et Biophysica Acta, (1973) 329/2 (221-232).

CODEN: BBACAQ

DOCUMENT TYPE: Journal

FILE SEGMENT: 037 Drug Literature Index

029 Clinical Biochemistry

025 Hematology 030 Pharmacology

LANGUAGE: English

AB The proteolytic action of human and bovine Factor Xa, bovine thrombin and bovine pancreatic trypsin on human Factor II at pH 7.5 and 25°C was monitored by sodium dodecylsulfate gel electrophoresis and thrombin assays. Purified human and bovine Factor Xa, and trypsin, were found to activate Factor II to thrombin. The conversion of Factor II to thrombin by either Factor Xa or trypsin was found to proceed through two thrombogenic intermediates. The reaction pathway appears to be sequential in that the Factor II (75000 daltons) is first cleaved to a 55000 dalton thrombogenic product (Intermediate 1) and a 25000 dalton non thrombogenic product (Fragment 1). Intermediate 1 is subsequently converted to an

inactive 37000 dalton thrombogenic protein (Intermediate 2) and a 16000 dalton protein (Fragment 2). Intermediate 2 is finally converted to an active 37000 dalton thrombin (α thrombin). Purified bovine thrombin readily converted Factor II to Intermediate 1 and Fragment 1, but possessed little capacity to catalyze subsequent cleavages to produce active thrombin. The ability of thrombin to cleave Factor II was entirely obviated in the presence of hirudin. Under the conditions of the incubation, the maximum thrombin yield obtainable by Factor Xa or trypsin activation was 50% when compared to the two stage potential thrombin.

=> file uspatfull europatfull pctfull
COST IN U.S. DOLLARS

FULL ESTIMATED COST

SINCE FILE TOTAL ENTRY SESSION 40.07 40.28

FILE 'USPATFULL' ENTERED AT 12:04:20 ON 17 JUN 2003 CA INDEXING COPYRIGHT (C) 2003 AMERICAN CHEMICAL SOCIETY (ACS)

FILE 'EUROPATFULL' ENTERED AT 12:04:20 ON 17 JUN 2003 COPYRIGHT (c) 2003 WILA Verlag Muenchen (WILA)

FILE 'PCTFULL' ENTERED AT 12:04:20 ON 17 JUN 2003 COPYRIGHT (C) 2003 Univentio

=> s 37000 adj (Da or dalton)
MISSING OPERATOR 'ADJ (DA'
The search profile that was entered contains terms or
nested terms that are not separated by a logical operator.

=> s 37000 (s) (Da or dalton) L7 29 37000 (S) (DA OR DALTON)

=> s 17 and EGFR

L8 0 L7 AND EGFR

=> d ibib abs kwic 19

·'ABS' IS NOT A VALID FORMAT FOR FILE 'EUROPATFULL'

The following are valid formats:

MAX ------ AN, ED, UP, EW, FS, TI, TIDE, TIFR, IN, PA, PAN, AG, AGN, OS, SO, DT, LA, DS, PIT, PI, OD, AI, PRAI, RLI, NTE, REP, REN, IC (ICM, ICS), ICA, ICI, CM, FA, GIS, PGC, CLMN, AB, ABDE, ABFR, DETD, DETDDE, DETDFR, CLMDE, CLMFR

MAXG ----- MAX plus GI MAX.OS ---- MAX, OS only MAXG.OS ---- MAX.OS plus GI MAX.PS ---- MAX, PS only

IMAX ----- MAX, indented with text labels

IMAX.OS ---- MAX, indented with text labels, OS only IMAX.PS ---- MAX, indented with text labels, PS only

ALL ------ AN, ED, UP, EW, FS, TI, IN, PA, PAN, AG, AGN, OS, SO, DT, LA, DS, PIT, PI, OD, AI, PRAI, RLI, NTE, REP, REN, IC (ICM, ICS), ICA, ICI, CM, FA, GIS, PGC, CLMN, AB*, DETD*, CLM* (* German or French text if English text is not available)

ALLG ----- ALL plus GI

```
ALL.OS ---- ALL, OS only
ALLG.OS ---- ALL.OS plus GI
ALL.PS ---- ALL, PS only IALL ----- ALL, indented with text labels
IALLG ----- IALL plus GI
IALL.OS ---- ALL, indented with text labels, OS only
IALLG.OS --- IALL.OS plus GI
IALL.PS ---- ALL, indented with text labels, PS only
ALLDE ----- AN, ED, UP, EW, FS, TIDE, IN, PA, PAN, AG, AGN, OS, SO, DT,
             LA, DS, PIT, PI, OD, AI, PRAI, RLI, NTE, REP, REN, IC (ICM,
             ICS), ICA, ICI, CM, FA, GIS, PGC, CLMN, ABDE* , DETDDE*, CLMDE*
             (* English or French text if German text is not available)
ALLGDE ---- ALLDE plus GI
ALLDE.OS --- ALLDE, OS only
ALLGDE.OS -- ALLDE.OS plus GI
ALLDE.PS --- ALLDE, PS only
ALLFR ----- AN, ED, UP, EW, FS, TIFR, IN, PA, PAN, AG, AGN, OS, SO, DT,
            LA, DS, PIT, PI, OD, AI, PRAI, RLI, NTE, REP, REN, IC (ICM,
             ICS), ICA, ICI, CM, FA, GIS, PGC, CLMN, ABFR* , DETDFR*, CLMFR*
             (* English or German text if French text is not available)
ALLGFR ---- ALLFR plus GI
ALLFR.OS --- ALLFR, OS only
ALLGFR.OS -- ALLFR.OS plus GI
ALLFR.PS --- ALLFR, PS only
BRIEF ----- AN, ED, UP, EW, FS, TI, IN, PA, PAN, AG, AGN, OS, SO, DT,
            LA, DS, PIT, PI, OD, AI, PRAI, RLI, NTE, REP, REN, IC (ICM,
            ICS), ICA, ICI, CM, FA, GIS, PGC, CLMN, AB*, MCLM*
             (* German or French text if English text is not available)
BRIEFG ---- BRIEF plus GI
BRIEF.OS --- BRIEF, OS only
BRIEFG.OS -- BRIEF.OS plus GI
BRIEF.PS --- BRIEF, PS only
IBRIEF ---- BRIEF, indented with text labels
IBRIEFG ---- IBRIEF plus GI
IBRIEF.OS -- BRIEF, indented with text labels, OS only
IBRIEFG.OS - IBRIEF.OS plus GI
IBRIEF.PS -- BRIEF, indented with text labels, PS only
BIB ----- AN, ED, UP, EW, FS, TI, TIDE, TIFR, IN, PA, PAN, AG, AGN,
            OS, SO, DT, LA, DS, PIT, PI, OD, AI, PRAI, RLI, NTE, REP, REN
BIB.OS ---- BIB, OS only
BIB.PS ---- BIB, PS only
IBIB ----- BIB, indented with text labels
IBIB.OS ---- BIB, indented with text labels, OS only
IBIB.PS ---- BIB, indented with text labels, PS only
BIBU ----- BIB, with German headers
BIBU.OS ---- BIB, with German headers, OS only
BIBU.PS ---- BIB, with German headers, PS only
STD ----- AN, ED, UP, EW, FS, TI, TIDE, TIFR, IN, PA, SO, DS, PIT, PI,
             OD, AI, PRAI, RLI, NTE, REP, REN, IC (ICM, ICS), ICA, ICI
STD.OS ---- STD, OS only
STD.PS ---- STD, PS only
ISTD ----- STD, indented with text labels
ISTD.OS ---- STD, indented with text labels, OS only
ISTD.PS ---- STD, indented with text labels, PS only
STDU ----- STD, with German headers
STDU.OS ---- STD, with German headers, OS only
STDU.PS ---- STD, with German headers, PS only
IND ----- ED, UP, EW, FS, IC (ICM, ICS), ICA, ICI
```

IND.OS ---- IND, OS only

IND.PS ---- IND, PS only

TRI ----- TI, TIDE, TIFR, IC (ICM, ICS), ICA, ICI, CLMN, PGC, FA, GIS

TRI.OS ---- TRI, OS only TRI.PS ---- TRI, PS only

TX ----- DETD, CLM TX.OS ----- TX, OS only TX.PS ----- TX, PS only TXDE ----- DETDDE, CLMDE TXDE.OS ---- TXDE, OS only TXDE . PS ---- TXDE, PS only

TXFR ----- DETDFR, CLMFR TXFR.OS ---- TXFR, OS only

TXFR.PS ---- TXFR, PS only

To display a particular field or fields, enter the display field codes. For a list of display field codes, enter 'HELP DFIELDS' at an arrow prompt (=>). Examples of formats include: 'TI'; 'AN,TI,AU'; 'BIB,CT'; 'TI,CT'.

The order of the terms in the formats is not important, but information will be displayed in the same order as the format specification.

The same formats may be used with the DISPLAY ACC command to display the record for a specified accession number.

ENTER DISPLAY FORMAT (STD): ibib

L9 ANSWER 1 OF 1 EUROPATFULL COPYRIGHT 2003 WILA

PATENT APPLICATION - PATENTANMELDUNG - DEMANDE DE BREVET

ACCESSION NUMBER:

TITLE:

1.176200 EUROPATFULL EW 200205 FS OS

Use of polyeptides or their encoding nucleic acids for the diagnosis or treatment of skin diseases or wound healing and their use in indentifying pharmacologically

acitve substances.

Verwendung von Polypeptiden oder diese kodierende Nukleinsaeuren zur Diagnose oder Behandlung von

Hauterkrankung oder Wundheilung sowie ihre Verwendung zur Indentifizierung von pharmakologisch aktiven

Substanzen.

Utilisation des polypeptides ou leurs acides nucleiques pour la diagnose ou traitement des maladies de la peau

ou de la cicatrisation de blessures et leurs utilisations pour l'identification des substances

pharmacologiquement actives.

INVENTOR(S): Halle, Joern-Peter, Nonnenwaldstr. 8c, 82377 Penzberg,

DE;

Goppelt, Andreas, Blutenburgstr. 30, 80636 Muenchen, DE;

Regenbogen, Johannes, Planegger Str. 1, 82152

Martinsried, DE;

Werner, Sabine, Titlisstr. 37, 8032 Zuerich, CH; Wolf, Eckhard, Wilhelmshof 10, 85764 Oberschleissheim,

PATENT ASSIGNEE(S): Switch Biotech Aktiengesellschaft, Fraunhoferstrasse 10,

82152 Martinsried, DE

PATENT ASSIGNEE NO:

3166290

AGENT:

Boesl, Raphael, Dr. rer. nat., Dipl.-Chem., Patent- und Rechtsanwaelte Bardehle . Pagenberg . Dost . Altenburg . Geissler . Isenbruck Galileiplatz 1, 81679 Muenchen, DE

AGENT NUMBER:

74943

OTHER SOURCE:

BEPA2002011 EP 1176200 A2 0108

SOURCE:

Wila-EPZ-2002-H05-T1a

DOCUMENT TYPE:

Patent

LANGUAGE:

Anmeldung in Deutsch; Veroeffentlichung in Deutsch

R AT; R BE; R CH; R CY; R DE; R DK; R ES; R FI; R FR; R DESIGNATED STATES:

GB; R GR; R IE; R IT; R LI; R LU; R MC; R NL; R PT; R

SE; R TR; R AL; R LT; R LV; R MK; R RO; R SI

PATENT INFO.PUB.TYPE: EPA2 EUROPAEISCHE PATENTANMELDUNG

PATENT INFORMATION:

	PATENT NO		KIND	DATE	
	EP	1176200	A2	20020130	
'OFFENLEGUNGS' DATE:				20020130	
APPLICATION INFO.:	ΕP	2001-112963		20010607	
PRIORITY APPLN. INFO.:	DΕ	2000-10030149		20000620	
•	US	2000-222081		20000801	

---Logging off of STN---

Executing the logoff script...

=> LOG Y.

COST IN U.S. DOLLARS SINCE FILE TOTAL ENTRY SESSION FULL ESTIMATED COST 8.52 48.80

STN INTERNATIONAL LOGOFF AT 12:07:56 ON 17 JUN 2003